

**Cross-linking ligation and sequencing of hybrids (qCLASH)
reveals an unpredicted miRNA targetome in melanoma
cells**

Kozar *et al.*,

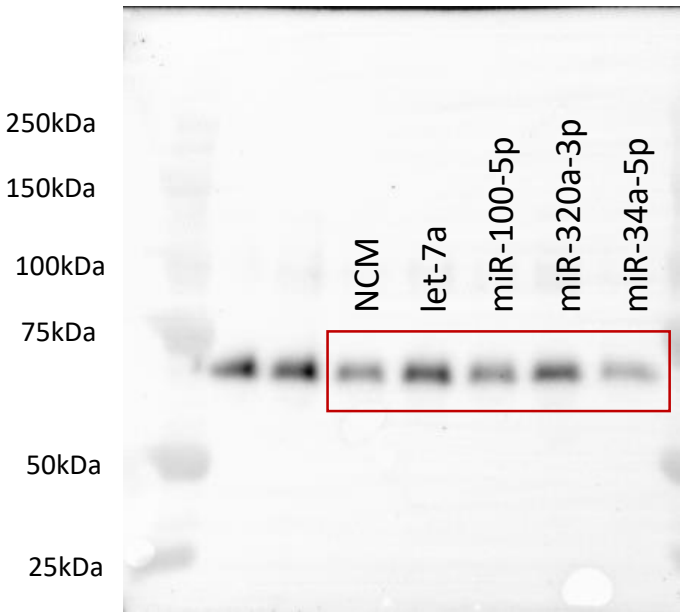
Supplements

Original western blot images corresponding to Figure 8B

*Parts of the blot that are shown in the manuscript are
highlighted by a red box*

*The original and non-annotated images are in the attached
zipped folder*

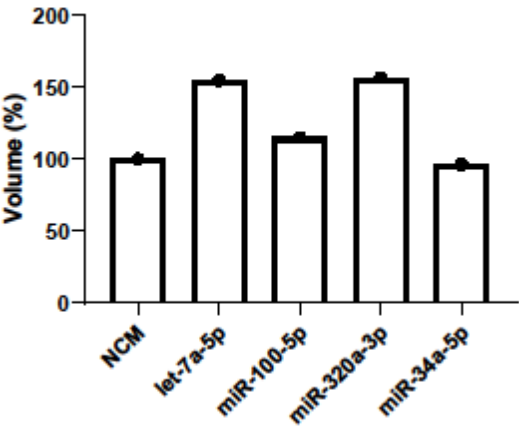
CD68



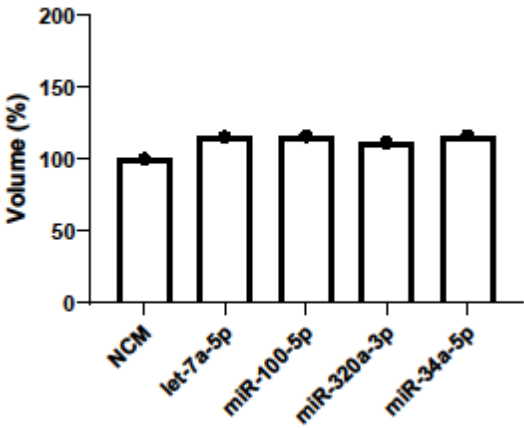
TUB



CD68



TUB

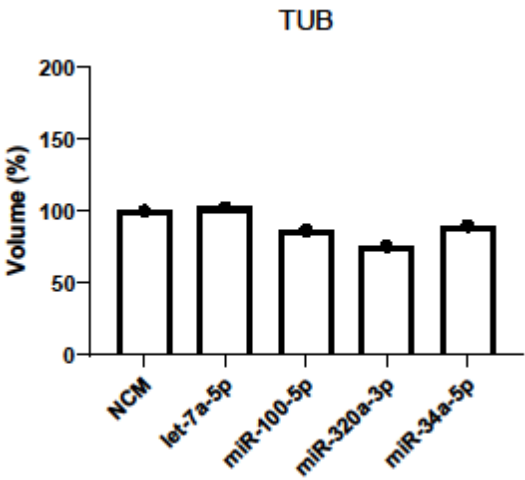
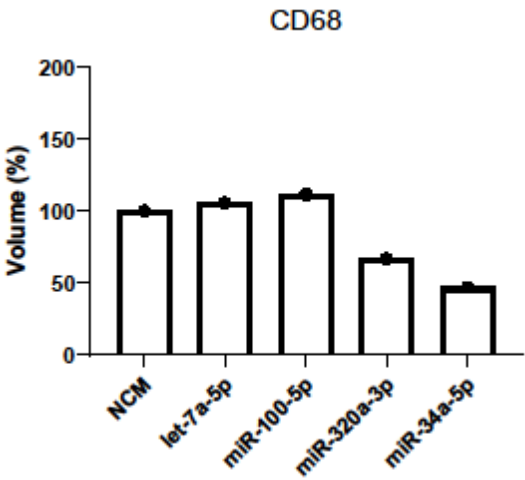
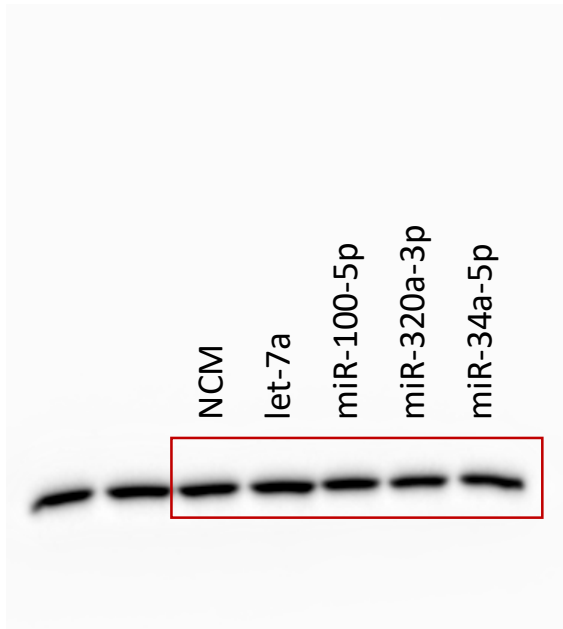
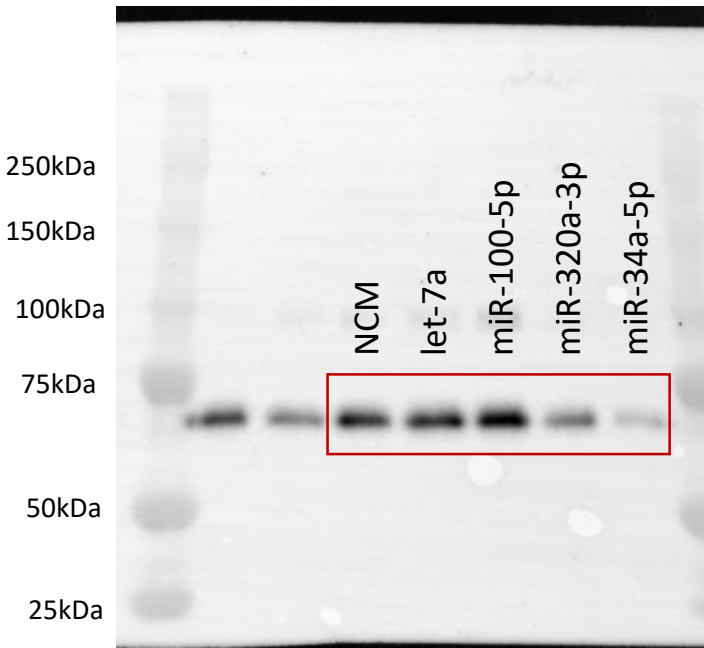


Protein bands have been quantified using the FusionCapt Advance software. The band intensity is expressed as the volume (%) relative to the intensity of the negative control mimic (NCM).

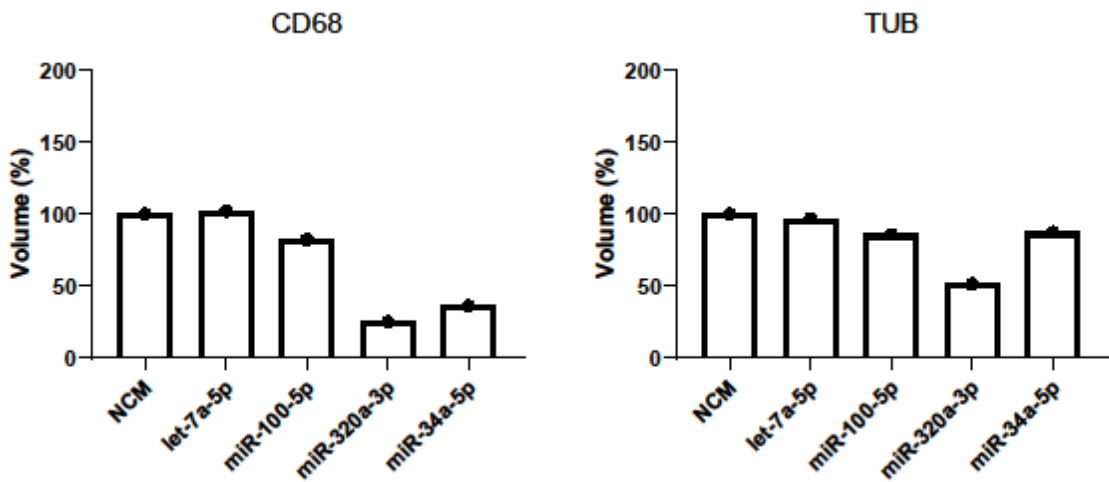
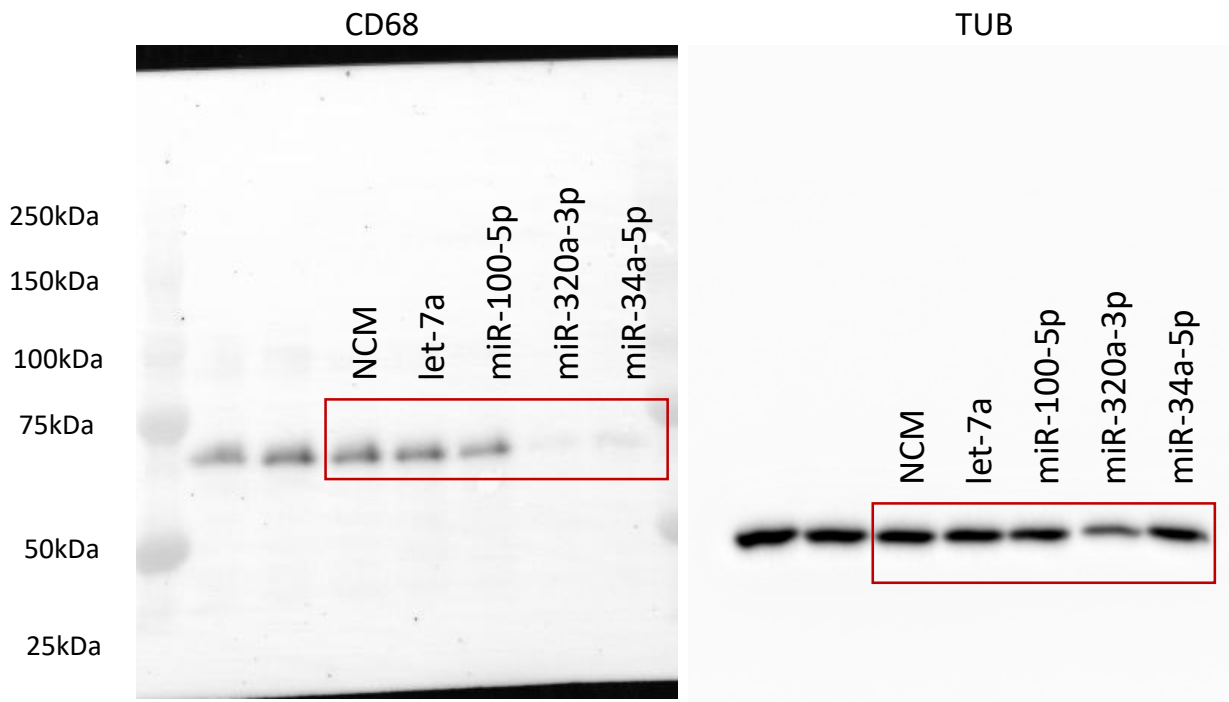
IGR37

CD68

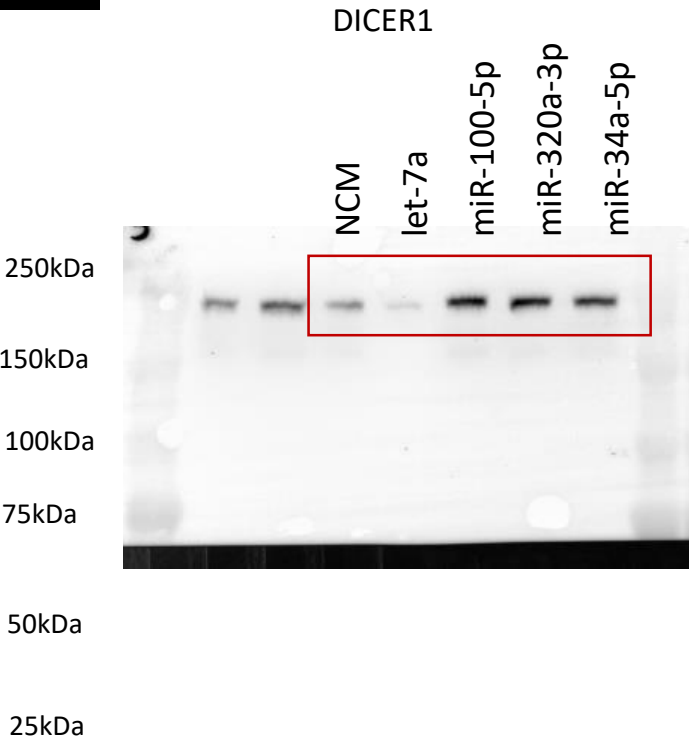
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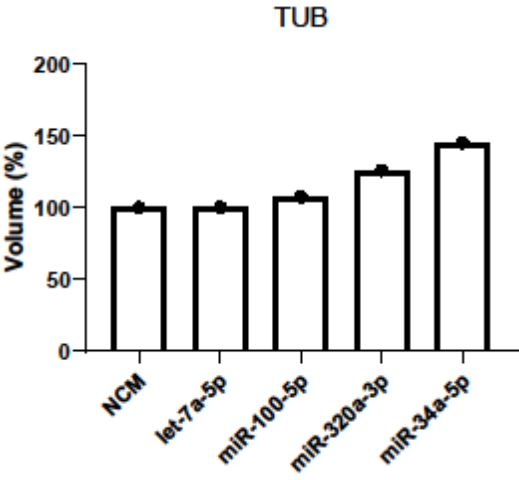
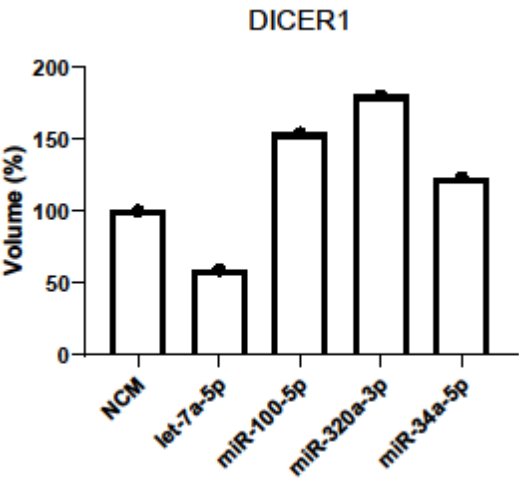
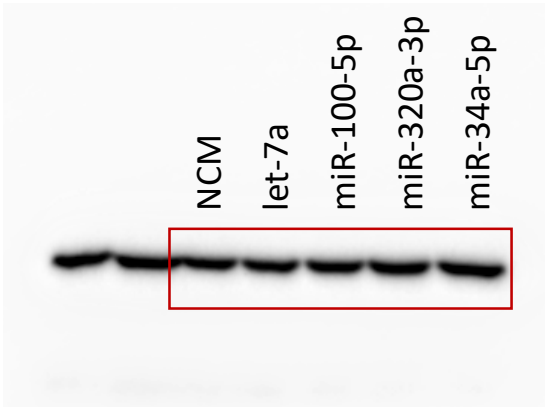
Protein bands have been quantified using the FusionCapt Advance software. The band intensity is expressed as the volume (%) relative to the intensity of the negative control mimic (NCM).



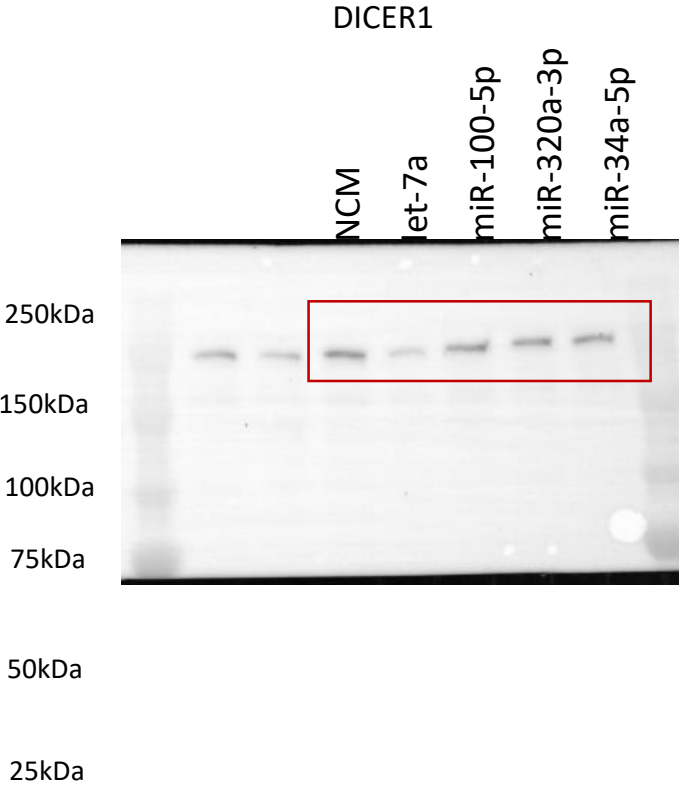
Protein bands have been quantified using the FusionCapt Advance software. The band intensity is expressed as the volume (%) relative to the intensity of the negative control mimic (NCM).



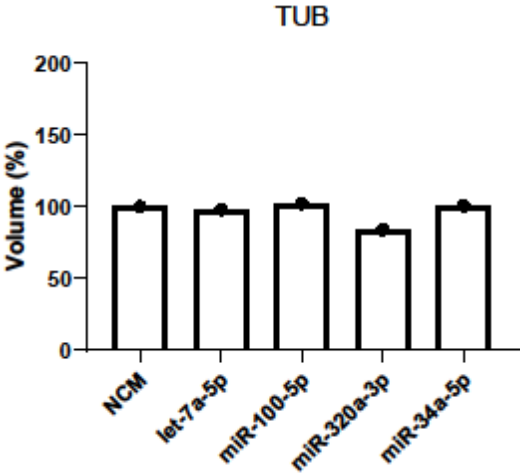
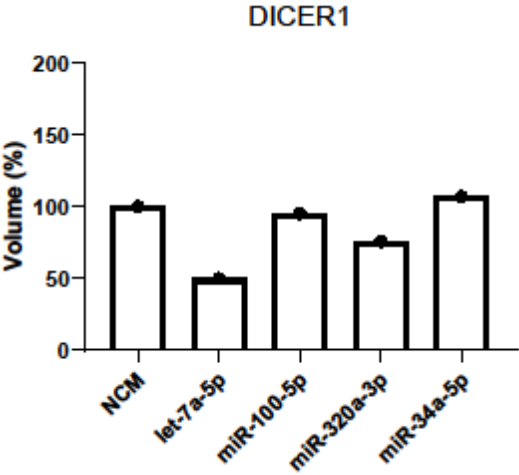
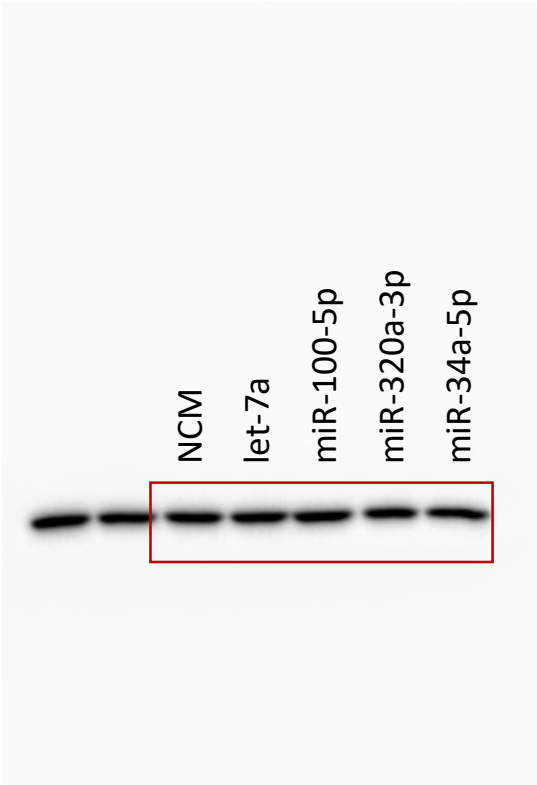
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Protein bands have been quantified using the FusionCapt Advance software. The band intensity is expressed as the volume (%) relative to the intensity of the negative control mimic (NCM).



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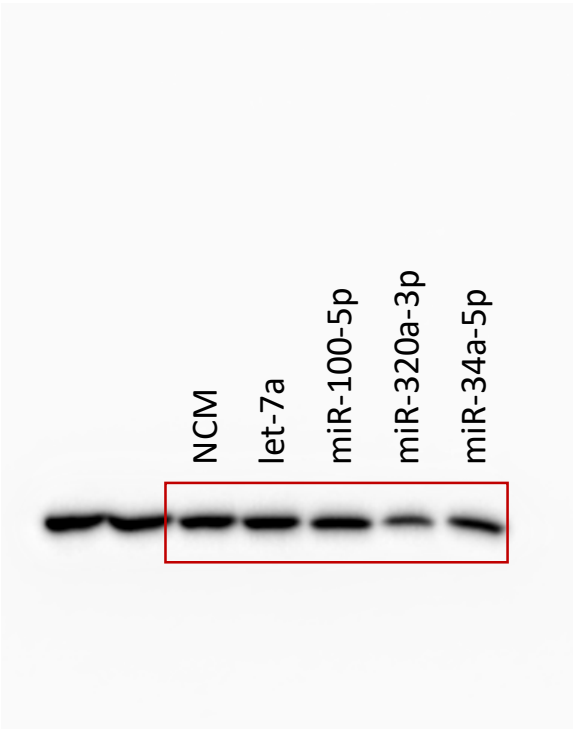
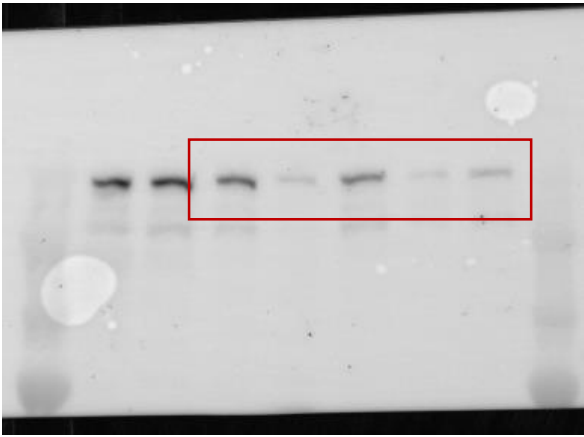
Protein bands have been quantified using the FusionCapt Advance software. The band intensity is expressed as the volume (%) relative to the intensity of the negative control mimic (NCM).

DICER1

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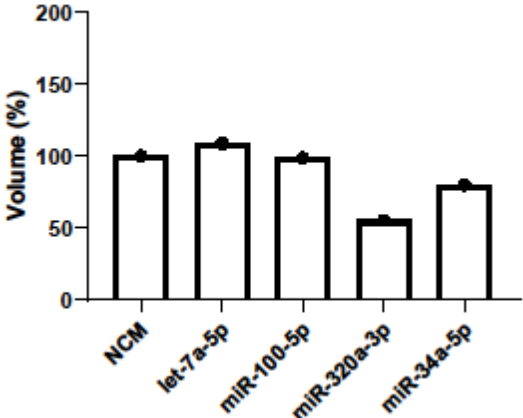
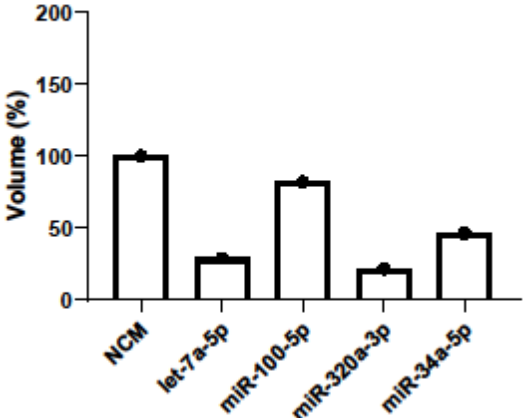
NCM
let-7a
miR-100-5p
miR-320a-3p
miR-34a-5p

250kDa
150kDa
100kDa
75kDa
50kDa
25kDa



DICER1

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Protein bands have been quantified using the FusionCapt Advance software. The band intensity is expressed as the volume (%) relative to the intensity of the negative control mimic (NCM).